Perper

## **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	_ <i>10   5</i> 07,928
Source:	P4/10
Date Processed by STIC:	6/10/05

# ENTERED



PCT

**RAW SEQUENCE LISTING**PATENT APPLICATION: **US/10/507,928**DATE: 06/10/2005
TIME: 10:08:51

Input Set : A:\Sequence Listing

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      5 <120> TITLE OF INVENTION: ADJUVANT
      7 <130> FILE REFERENCE: N.88232B GCW
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C--> 9 <141> CURRENT FILING DATE: 2004-09-17
      9 <160> NUMBER OF SEQ ID NOS: 12
     11 <170> SOFTWARE: PatentIn version 3.1
     13 <210> SEO ID NO: 1
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     15 <212> TYPE: PRT
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                                                                              120
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                                                                              180
     54 cttgaacggt ttgccgtgaa cccaggcctg ctggaaacat ctgagggatg tcgccagatc
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     56 ctggggcaat tgcagccatc cctccagacc gggagtgaag agctgaggtc cttgtataac
     58 acagtggcta ccctctactg cgtacaccag aggatcgaga ttaaggatac caaggaggcc
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     60 ttgqacaaaa ttgaggagga gcaaaacaag agcaagaaga aggcccagca ggcagctgct
                                                                              360
     62 gacactgggc atagcaacca ggtatcacag aactatccta ttgtccaaaa cattcagggc
                                                                              420
     64 cagatggttc atcaggccat cagccccgg acgctcaatg cctgggtgaa ggttgtcgaa
                                                                              480
     66 gagaaggeet ttteteetga ggttateeee atgtteteeg etttgagtga gggggeeaet
                                                                              540
                                                                              600
     68 cctcaggacc tcaatacaat gcttaatacc gtgggcggcc atcaggccgc catgcaaatg
     70 ttgaaggaga ctatcaacga ggaggcagcc gagtgggaca gagtgcatcc cgtccacgct
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DATE: 06/10/2005 TIME: 10:08:51

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/507,928

Input Set : A:\Sequence Listing

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                                                                         780
76 atctataaac ggtggatcat tctcggtctc aataaaattg ttagaatgta ctctccgaca
                                                                         840
78 tccatccttg acattagaca gggacccaaa gagcctttta gggattacgt cgaccggttt
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80 tataagaccc tgcgagcaga gcaggcctct caggaggtca aaaactggat gacggagaca
                                                                         960
82 ctcctggtac agaacgctaa ccccgactgc aaaacaatct tgaaggcact aggcccggct
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84 gccaccetgg aagagatgat gaccgcetgt cagggagtag gcggaccegg acacaaagce
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86 agagtgttgg ccgaagccat gagccaggtg acgaactccg caaccatcat gatgcagaga
                                                                        1140
                                                                        1200
88 gggaacttcc gcaatcagcg gaagatcgtg aagtgtttca attgcggcaa ggagggtcat
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90 accqcccqca actqtcqqqc ccctaggaag aaagggtgtt ggaagtgcgg caaggaggga
92 caccagatga aagactgtac agaacgacag gccaattttc ttggaaagat ttggccgagc
                                                                        1320
94 tacaagggga gacctggtaa tttcctgcaa agcaggcccg agcccaccgc ccccctgag
                                                                        1380
96 gaatcettca ggteeggagt ggagaceaea aegeeteeee aaaaaeagga aecaategae
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118
121 His Ile Val Trp Ala Ser Arg Glu Leu Glu Arg Phe Ala Val Asn Pro
122
125 Gly Leu Leu Glu Thr Ser Glu Gly Cys Arg Gln Ile Leu Gly Gln Leu
                            55
129 Gln Pro Ser Leu Gln Thr Gly Ser Glu Glu Leu Arg Ser Leu Tyr Asn
                        70
                                             75
133 Thr Val Ala Thr Leu Tyr Cys Val His Gln Arg Ile Glu Ile Lys Asp
                                         90
                    85
137 Thr Lys Glu Ala Leu Asp Lys Ile Glu Glu Glu Gln Asn Lys Ser Lys
138
                100
                                     105
141 Lys Lys Ala Gln Gln Ala Ala Ala Asp Thr Gly His Ser Asn Gln Val
                                120
                                                     125
            115
145 Ser Gln Asn Tyr Pro Ile Val Gln Asn Ile Gln Gly Gln Met Val His
                            135
        130
149 Gln Ala Ile Ser Pro Arq Thr Leu Asn Ala Trp Val Lys Val Val Glu
                                             155
150 145
                        150
153 Glu Lys Ala Phe Ser Pro Glu Val Ile Pro Met Phe Ser Ala Leu Ser
                                         170
154
                    165
157 Glu Gly Ala Thr Pro Gln Asp Leu Asn Thr Met Leu Asn Thr Val Gly
                                    185
161 Gly His Gln Ala Ala Met Gln Met Leu Lys Glu Thr Ile Asn Glu Glu
            195
                                200
165 Ala Ala Glu Trp Asp Arg Val His Pro Val His Ala Gly Pro Ile Ala
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TIME: 10:08:51

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/507,928

Input Set : A:\Sequence Listing

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173 Ser Thr Leu Gln Glu Gln Ile Gly Trp Met Thr Asn Asn Pro Pro Ile 174				-			-			-	_									
174				Thr	Leu	Gln	Glu	Gln	Ile	Glv	Trp	Met	Thr	Asn	Asn	Pro	Pro	Ile		
177    Pro Val Gly Glu Ile Tyr Lys Arg Trp Ile Ile Leu Gly Leu Asn Lys 178    260										-4	•									
178			Pro	Va 1	Glv	Glu		Tur	Lvs	Ara	Trn		Ile	Leu	Glv	Leu		Lvs		
181   11e Val Arg Met Tyr Ser Pro Thr Ser Ile Leu Asp Ile Arg Gln Gly 182   275   280   285   28			110	Val	C±y			- 1 -	_,_	9								-1-		
182			T10	Val	Ara		Tur	Sar	Pro	Thr		Tle	T.e.11	Asn	Tle		Gln	Glv		
185  Pro Lys Glu Pro Phe Arg Asp Tyr Val Asp Arg Phe Tyr Lys Thr Leu 186  290  295  300 189  Arg Ala Glu Gln Ala Ser Gln Glu Val Lys Asn Trp Met Thr Glu Thr 190 305  310  315  320 193  Leu Leu Val Gln Asn Ala Asn Pro Asp Cys Lys Thr Ile Leu Lys Ala 194  325  330  335 197  Leu Gly Pro Ala Ala Thr Leu Glu Glu Met Met Thr Ala Cys Gln Gly 198  340  345  350 201 Val Gly Gly Pro Gly His Lys Ala Arg Val Leu Ala Glu Ala Met Ser 202  355  360  365 205  Gln Val Thr Asn Ser Ala Thr Ile Met Met Gln Arg Gly Asn Phe Arg 206  370  375  380 209  Asn Gln Arg Lys Ile Val Lys Cys Phe Asn Cys Gly Lys Glu Gly His 210  385  390  395 213  Thr Ala Arg Asn Cys Arg Ala Pro Arg Lys Lys Gly Cys Trp Lys Cys 214  405  410  415 217  Gly Lys Glu Gly His Gln Met Lys Asp Cys Thr Glu Arg Gln Ala Asn 218  420  425 221  Phe Leu Gly Lys Ile Trp Pro Ser Tyr Lys Gly Arg Pro Gly Asn Phe 222  435  440 221  Phe Leu Gly Lys Ile Trp Pro Ser Tyr Lys Gly Arg Pro Gly Asn Phe 222  435  440 223  Lys Glu Leu Tyr Pro Leu Thr Ser Leu Arg Ser Leu Phe Gly Asn Asp 233  Lys Glu Leu Tyr Pro Leu Thr Ser Leu Arg Ser Leu Phe Gly Asn Asp 244  4213  ORGANISM: Artificial sequence 246  <220> FEAURE: 247  <221> LENGTH: 1515 243  <212> TYFE: DNA 244  <213> ORGANISM: Artificial sequence 246  <220> FEAURE: 247  <223> OTHER INFORMATION: nucleotide sequence of the pl7/24trNEF insert in 249  <400> SEQUENCE: 5 250  atgggtgoga gagggaaaga aaaatataaa ttaaaacata tagtatggga aaaaattcgg 60 252  ttaaggccag ggggaaagaa aaaatataaa ttaaaacata tagtatggga aagaagagag 120 254  ctaggacagt tcaaacacat ccttcagaca ggatcaagaacat cagaagagtt tattatataat 240			116	Val		ricc	+ y +	DCI	110		DCI	110	104	1102		9	02	0-1		
186			Dro	Tuc		Dro	Pho	Ara	Δen		17 a 1	Asn	Δra	Phe		T.vs	Thr	Len		
189 Arg Ala Glu Gln Ala Ser Gln Glu Val Lys Asn Trp Met Thr Glu Thr 190 305 310 310 315 320  193 Leu Leu Val Gln Asn Ala Asn Pro Asp Cys Lys Thr Ile Leu Lys Ala 194 325 330 335  197 Leu Gly Pro Ala Ala Thr Leu Glu Glu Met Met Thr Ala Cys Gln Gly 198 340 345 350  201 Val Gly Gly Pro Gly His Lys Ala Arg Val Leu Ala Glu Ala Met Ser 202 355 360 365  205 Gln Val Thr Asn Ser Ala Thr Ile Met Met Gln Arg Gly Asn Phe Arg 206 370 375 380  209 Asn Gln Arg Lys Ile Val Lys Cys Phe Asn Cys Gly Lys Glu Gly His 210 385 390 395 400  213 Thr Ala Arg Asn Cys Arg Ala Pro Arg Lys Lys Gly Cys Trp Lys Cys 214 405 415  217 Gly Lys Glu Gly His Gln Met Lys Asp Cys Thr Glu Arg Gln Ala Asn 218 420 425 435  221 Phe Leu Gly Lys Ile Trp Pro Ser Tyr Lys Gly Arg Pro Gly Asn Phe 222 435 440  223 Lys Glu Ser Arg Pro Glu Pro Thr Ala Pro Pro Glu Glu Ser Phe Arg 226 450 455 460  229 Ser Gly Val Glu Thr Thr Thr Pro Pro Gln Lys Gln Glu Pro Ile Asp 230 465 470 455  231 Lys Glu Leu Tyr Pro Leu Thr Ser Leu Arg Ser Leu Phe Gly Asn Asp 232 Pro Ser Ser Gln 233 Lys Glu Leu Tyr Pro Leu Thr Ser Leu Arg Ser Leu Phe Gly Asn Asp 246 <220> FEATURE: 247 <223> ORGANISM: Artificial sequence 246 <220> FEATURE: 247 <223> ORTHER INFORMATION: nucleotide sequence of the p17/24trNEF insert in p17/24trNEF1 249 <400> SEQUENCE: 5 250 atgggtgcag gaggaaagaa aaaatataaa ttaaaacata tagatagggag aaaaattcgg 60 252 ttaaaggccag ggggaaagaa aaaatataaa ttaaaacata tagataggc aagacaggaga 120 254 ctagaacagt tacaacacata cottcagaca ggatcaagaa aacatataaat 180 256 ctgggacagc tacaacacata cottcagaca ggatcaagaa aacatataaat 2440			FIO		GIU	FIO	1116	Ary		ı yı	vaı	пор	nra		- y -	Lys	1	LCu		
190 305			7\ ~~ ~		Clu	Cln	712	Sor		Glu	Val	Twe	Δen		Mot	Thr	Glu	Thr		
193 Leu Leu Val Gln Asn Ala Asn Pro Asp Cys Lys Thr Ile Leu Lys Ala 194 325 330 335 197 Leu Gly Pro Ala Ala Thr Leu Glu Glu Met Met Thr Ala Cys Gln Gly 198 340 345 350 201 Val Gly Gly Pro Gly His Lys Ala Ary Val Leu Ala Glu Ala Met Ser 202 355 360 365 205 Gln Val Thr Asn Ser Ala Thr Ile Met Met Gln Arg Gly Asn Phe Arg 206 370 375 380 209 Asn Gln Arg Lys Ile Val Lys Cys Phe Asn Cys Gly Lys Glu Gly His 210 385 390 395 213 Thr Ala Arg Asn Cys Arg Ala Pro Arg Lys Lys Gly Cys Trp Lys Cys 214 405 410 415 217 Gly Lys Glu Gly His Gln Met Lys Asp Cys Thr Glu Arg Gln Ala Asn 218 420 425 430 221 Phe Leu Gly Lys Ile Trp Pro Ser Tyr Lys Gly Arg Pro Gly Asn Phe 222 435 450 450 229 Ser Gly Val Glu Thr Thr Thr Pro Pro Gln Lys Gly Glu Ser Phe Arg 230 465 470 233 Lys Glu Leu Tyr Pro Leu Thr Ser Leu Arg Ser Leu Phe Gly Asn Asp 234 4210 SEQ Ile Leu Thr Ser Leu Arg Ser Leu Phe Gly Asn Asp 235 400 241 <210> SEQ ID NO: 5 242 <211> LENGTH: 1515 243 <212> TYPE: DNA 244 <223> OTHER INFORMATION: nucleotide sequence of the p17/24trNEF insert in 247 <220> Ser Glyggaaagaa aaaatataaa ttaaaacata tagtaggga aaaaattcgg 60 252 ttaaggccag ggggaaagaa aaaatataaa ttaaaacata tagtaggga aagacaggaga 120 254 ctagaacgat tcacaaccatc cottcagaca ggatcagaag aacattagat cattaataat 240			_	Ата	GIU	GIII			GIII	GIU	vaı	ъйз		пр	Het	1111	Olu			
194				T	17-1	C1-			Λον	Dro	7 an	Cvc		Thr	Tla	LOU	Lve			
197 Leu Gly Pro Ala Ala Thr Leu Glu Glu Met Met Thr Ala Cys Gln Gly 198 340 345 350 201 Val Gly Gly Pro Gly His Lys Ala Arg Val Leu Ala Glu Ala Met Ser 202 355 360 365 205 Gln Val Thr Asn Ser Ala Thr Ile Met Met Gln Arg Gly Asn Phe Arg 206 370 375 380 209 Asn Gln Arg Lys Ile Val Lys Cys Phe Asn Cys Gly Lys Glu Gly His 210 385 390 395 400 213 Thr Ala Arg Asn Cys Arg Ala Pro Arg Lys Lys Gly Cys Trp Lys Cys 214 405 410 415 217 Gly Lys Glu Gly His Gln Met Lys Asp Cys Thr Glu Arg Gln Ala Asn 218 420 425 400 221 Phe Leu Gly Lys Ile Trp Pro Ser Tyr Lys Gly Arg Pro Gly Asn Phe 222 435 400 222 Pro Glu Ser Arg Pro Glu Pro Thr Ala Pro Pro Glu Glu Ser Phe Arg 226 450 455 460 229 Ser Gly Val Glu Thr Thr Thr Pro Pro Gln Lys Gln Glu Pro Ile Asp 230 465 470 233 Lys Glu Leu Tyr Pro Leu Thr Ser Leu Arg Ser Leu Phe Gly Asn Asp 234 485 490 237 Pro Ser Ser Gln 238 500 241 <210> SEQ ID NO: 5 242 <211> LENGTH: 1515 243 <212> TYPE: DNA 244 <213> ORGANISM: Artificial sequence 246 <220> FEATURE: 247 <223> OTHER INFORMATION: nucleotide sequence of the pl7/24trNEF insert in pl7/24trNEF1 249 <400> SEQUENCE: 5 250 atgggtgcga gagcgtcagt attaagcggg ggagaattag atcgatggga aaaaattcgg 60 252 ttaaggccag ggggaaagaa aaaatataaa ttaaaacata tagtatgggc aagacagtgag 120 256 ctaggacagt tcgcagttaa tcctggcctg ttagaacacat cagaaggtg attatataat 240			ьeu	ьeu	Val	GIII		Ala	ASII	PIO	ASP		гуэ	1111	116	neu		ALG		
198				<b>01</b>	D	7.1.		m\	T	C1	C1		Mot	mb ~	717	C		C1,,		
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206			<b>~</b> 1				<b>G</b>	70.7 -	m1			M-4	C1-	7		7 ~~	Dho	7.~~		
209 Asn Gln Arg Lys Ile Val Lys Cys Phe Asn Cys Gly Lys Glu Gly His 210 385 390 395 400 213 Thr Ala Arg Asn Cys Arg Ala Pro Arg Lys Lys Gly Cys Trp Lys Cys 214 405 415 217 Gly Lys Glu Gly His Gln Met Lys Asp Cys Thr Glu Arg Gln Ala Asn 218 420 425 430 221 Phe Leu Gly Lys Ile Trp Pro Ser Tyr Lys Gly Arg Pro Gly Asn Phe 222 435 440 225 Leu Gln Ser Arg Pro Glu Pro Thr Ala Pro Pro Glu Glu Ser Phe Arg 226 450 455 229 Ser Gly Val Glu Thr Thr Thr Pro Pro Gln Lys Gln Glu Pro Ile Asp 230 465 470 475 480 231 Lys Glu Leu Tyr Pro Leu Thr Ser Leu Arg Ser Leu Phe Gly Asn Asp 234 485 490 495 237 Pro Ser Ser Gln 238 500 241 <210> SEQ ID NO: 5 242 <211> LENGTH: 1515 243 <212> TYPE: DNA 244 <213> ORGANISM: Artificial sequence 246 <220> FEATURE: 247 <223> OTHER INFORMATION: nucleotide sequence of the p17/24trNEF insert in p17/24trNEF1 249 <400> SEQUENCE: 5 250 atgggtgcga gagcgtcagt attaagcggg ggagaattag atcgatggga aaaaattcgg 60 252 ttaagaccgat tcgcagttaa tcctggcctg ttagaacaat tagtatgggc aagcaggag 120 254 ctagaacgat tcgcagttaa tcctggcctg ttagaacaat tagtattgagc agcaggag 120 254 ctagaacgat tcgcagttaa tccttggcctg ttagaacaat atgattagatc attatataat 240			GIn		Thr	Asn	Ser			TTE	Met	Met	GIII		СТУ	ASII	Pne	ALG		
210 385			_		_		- 1			<b>a</b>	D1	7	<b>0</b>		T	C1	C1	u.i.a		
213 Thr Ala Arg Asn Cys Arg Ala Pro Arg Lys Lys Gly Cys Trp Lys Cys 214				GIN	Arg	гàг	тте		гаг	Cys	Pne	ASII		GIY	гĀЗ	GIU	Gry			
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218				_	- 1	~1		<b>~</b> 1	24.1		7		m\	C1	7)	C1-		7.00		
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222			51	<b>.</b>	<b>61</b>		~1.	m	D	0		T	C1	7)	Dwa		7.00	Dho		
225 Leu Gln Ser Arg Pro Glu Pro Thr Ala Pro Pro Glu Glu Ser Phe Arg 226	•		Phe	ьeu		ьys	тте	Trp	Pro		Tyr	гуѕ	СТУ	Arg		СТУ	ASII	rne		
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250 atgggtgcga gagcgtcagt attaagcggg ggagaattag atcgatggga aaaaattcgg 60 252 ttaaggccag ggggaaagaa aaaatataaa ttaaaacata tagtatgggc aagcagggag 120 254 ctagaacgat tcgcagttaa tcctggcctg ttagaaacat cagaaggctg tagacaaata 180 256 ctgggacagc tacaaccatc ccttcagaca ggatcagaag aacttagatc attatataat 240	p17/2			<b>.</b> -			_													
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256 ctgggacagc tacaaccatc ccttcagaca ggatcagaag aacttagatc attatataat 240																				
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258 acagtagcaa ccctctattg tgtgcatcaa aggatagaga taaaagacac caaggaagct 300		256	ctg	ggac	agc ·	taca	acca	tc c	cttc	agac	a gg	atca	gaag	aac	ttag	atc	atta	tataat		
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TIME: 10:08:51

### RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/507,928

Input Set : A:\Sequence Listing

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     264 caaatqqtac atcaqqccat atcacctaqa actttaaatq catqqqtaaa aqtaqtagaa
                                                                               540
     266 gagaaggett teageecaga agtgataece atgtttteag cattateaga aggageeace
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     268 ccacaagatt taaacaccat qctaaacaca gtggggggac atcaagcagc catgcaaatg
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     270 ttaaaagaga ccatcaatga qgaagctgca gaatgggata gagtgcatcc agtgcatgca
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     272 gggcctattg caccaggcca gatgagagaa ccaaggggaa gtgacatagc aggaactact
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     274 agtaccette aggaacaaat aggatggatg acaaataate cacctateee agtaggagaa
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     278 agcattctgg acataagaca aggaccaaaa gaacccttta gagactatgt agaccggttc
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     286 agagttttgg tgggttttcc agtcacacct caggtacctt taagaccaat gacttacaag
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     288 gcagctgtag atcttagcca ctttttaaaa gaaaaggggg gactggaagg gctaattcac
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     296 agcttgttac accctgtgag cctgcatggg atggatgacc cggagagaga agtgttagag
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     317 Glu Lys Ile Arg Leu Arg Pro Gly Gly Lys Lys Lys Tyr Lys Leu Lys
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                                     40
     325 Gly Leu Leu Glu Thr Ser Glu Gly Cys Arg Gln Ile Leu Gly Gln Leu
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     329 Gln Pro Ser Leu Gln Thr Gly Ser Glu Glu Leu Arg Ser Leu Tyr Asn
                             70
                                                 75
     333 Thr Val Ala Thr Leu Tyr Cys Val His Gln Arg Ile Glu Ile Lys Asp
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     337 Thr Lys Glu Ala Leu Asp Lys Ile Glu Glu Glu Gln Asn Lys Ser Lys
     338
     341 Lys Lys Ala Gln Gln Ala Ala Ala Asp Thr Gly His Ser Asn Gln Val
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     345 Ser Gln Asn Tyr Pro Ile Val Gln Asn Ile Gln Gly Gln Met Val His
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     349 Gln Ala Ile Ser Pro Arg Thr Leu Asn Ala Trp Val Lys Val Val Glu
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### RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/507,928 TIME: 10:08:51

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                                   200
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    365 Ala Ala Glu Trp Asp Arg Val His Pro Val His Ala Gly Pro Ile Ala
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    369 Pro Gly Gln Met Arg Glu Pro Arg Gly Ser Asp Ile Ala Gly Thr Thr
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                            230
    373 Ser Thr Leu Gln Glu Gln Ile Gly Trp Met Thr Asn Asn Pro Pro Ile
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                                            250
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                                        265
    381 Ile Val Arg Met Tyr Ser Pro Thr Ser Ile Leu Asp Ile Arg Gln Gly
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                                    280
    385 Pro Lys Glu Pro Phe Arg Asp Tyr Val Asp Arg Phe Tyr Lys Thr Leu
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                                                    300
    389 Arg Ala Glu Gln Ala Ser Gln Glu Val Lys Asn Trp Met Thr Glu Thr
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    393 Leu Leu Val Gln Asn Ala Asn Pro Asp Cys Lys Thr Ile Leu Lys Ala
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     397 Leu Gly Pro Ala Ala Thr Leu Glu Glu Met Met Thr Ala Cys Gln Gly
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    401 Val Gly Gly Pro Gly His Lys Ala Arg Val Leu Val Gly Phe Pro Val
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    405 Thr Pro Gln Val Pro Leu Arg Pro Met Thr Tyr Lys Ala Ala Val Asp
                                375
     409 Leu Ser His Phe Leu Lys Glu Lys Gly Leu Glu Gly Leu Ile His
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     413 Ser Gln Arg Arg Gln Asp Ile Leu Asp Leu Trp Ile Tyr His Thr Gln
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                                            410
     417 Gly Tyr Phe Pro Asp Trp Gln Asn Tyr Thr Pro Gly Pro Gly Val Arg
                                        425
     421 Tyr Pro Leu Thr Phe Gly Trp Cys Tyr Lys Leu Val Pro Val Glu Pro
               435
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                                                        445
     425 Asp Lys Val Glu Glu Ala Asn Lys Gly Glu Asn Thr Ser Leu Leu His
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                                                    460
    429 Pro Val Ser Leu His Gly Met Asp Asp Pro Glu Arg Glu Val Leu Glu
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PATENT APPLICATION: US/10/507,928

DATE: 06/10/2005 TIME: 10:08:52

Input Set : A:\Sequence Listing

Output Set: N:\CRF4\06102005\J507928.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date